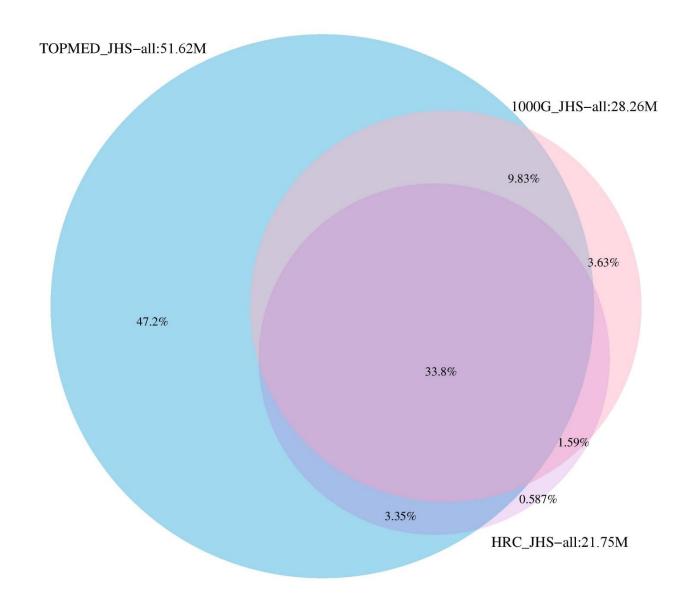
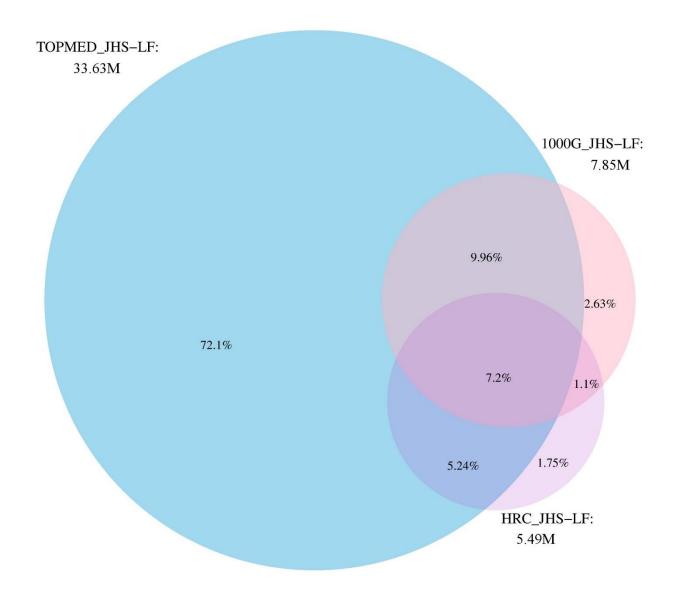
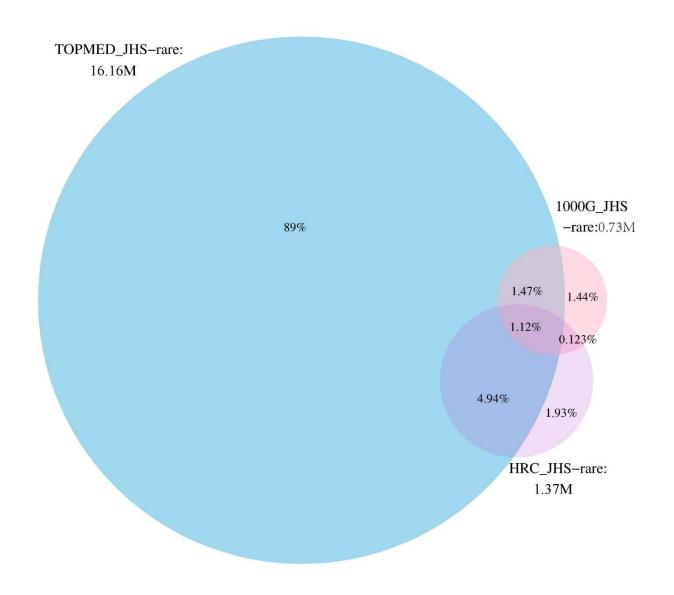
A. Comparison of all well-imputed variants included in results from TOPMed (TOPMed freeze 5b), 1000G (the 1000 Genomes Phase 3), and HRC (the Haplotype Reference Consortium) in the Jackson Heart Study (JHS).



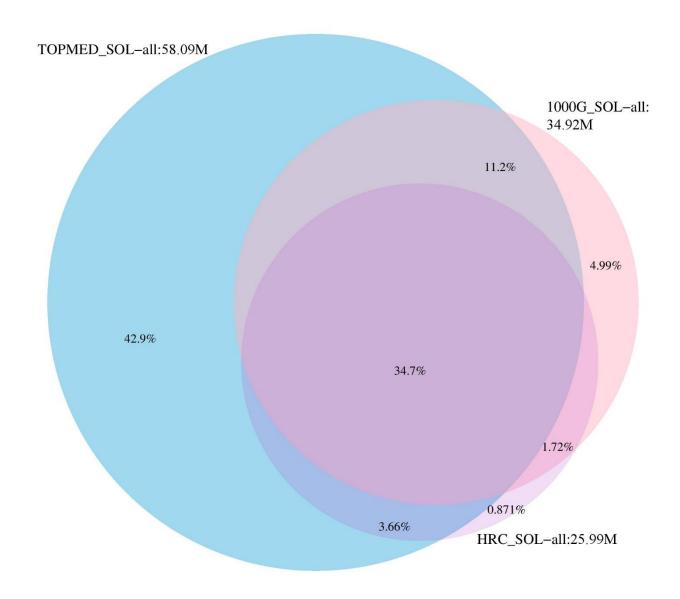
B. Comparison of all well-imputed variants with minor allele frequency < 0.5% included in results from TOPMed (TOPMed freeze 5b), 1000G (the 1000 Genomes Phase 3), and HRC (the Haplotype Reference Consortium) in the Jackson Heart Study (JHS).</p>



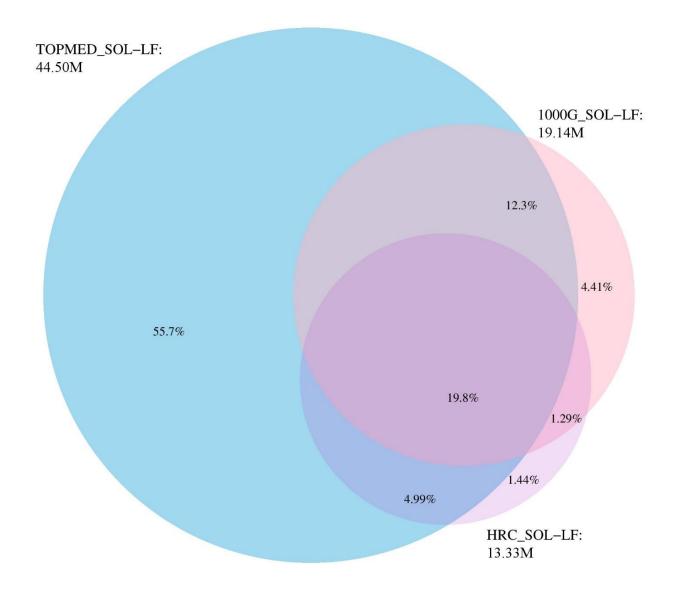
C. Comparison of all well-imputed variants with minor allele frequency < 0.05% included in results from TOPMed (TOPMed freeze 5b), 1000G (the 1000 Genomes Phase 3), and HRC (the Haplotype Reference Consortium) in the Jackson Heart Study (JHS).</p>



D. Comparison of all well-imputed variants included in results from TOPMed (TOPMed freeze 5b), 1000G (the 1000 Genomes Phase 3), and HRC (the Haplotype Reference Consortium) in the Hispanic Community Health Study/Study of Latinos (HCHS/SOL).



E. Comparison of all well-imputed variants with minor allele frequency < 0.5% included in results from TOPMed (TOPMed freeze 5b), 1000G (the 1000 Genomes Phase 3), and HRC (the Haplotype Reference Consortium) in the Hispanic Community Health Study/Study of Latinos (HCHS/SOL).



F. Comparison of all well-imputed variants with minor allele frequency < 0.05% included in results from TOPMed (TOPMed freeze 5b), 1000G (the 1000 Genomes Phase 3), and HRC (the Haplotype Reference Consortium) in the Hispanic Community Health Study/Study of Latinos (HCHS/SOL).

